



KAT6B gene

lysine acetyltransferase 6B

Normal Function

The *KAT6B* gene provides instructions for making a type of enzyme called a histone acetyltransferase. These enzymes modify histones, which are structural proteins that attach (bind) to DNA and give chromosomes their shape. By adding a small molecule called an acetyl group to particular locations on histones, histone acetyltransferases control the activity of certain genes.

Little is known about the function of the histone acetyltransferase produced from the *KAT6B* gene. It is active in cells and tissues throughout the body, where it interacts with many other proteins. It appears to regulate genes that are important for early development, including development of the skeleton and nervous system.

Health Conditions Related to Genetic Changes

genitopatellar syndrome

At least eight mutations in the *KAT6B* gene have been identified in people with genitopatellar syndrome, a rare condition characterized by genital abnormalities, missing or underdeveloped kneecaps (patellae), intellectual disability, and abnormalities affecting other parts of the body. The mutations that cause genitopatellar syndrome occur near the end of the *KAT6B* gene in a region known as exon 18. These mutations lead to the production of a shortened histone acetyltransferase enzyme. Researchers suspect that the shortened enzyme may function differently than the full-length version, altering the regulation of various genes during early development. Because the altered enzyme takes on a different function, these mutations are described as "gain-of-function." However, it is unclear how these changes lead to the specific features of genitopatellar syndrome.

Ohdo syndrome, Say-Barber-Biesecker-Young-Simpson variant

More than 10 mutations in the *KAT6B* gene have been found to cause the Say-Barber-Biesecker-Young-Simpson (SBBYS) variant of Ohdo syndrome. This condition has signs and symptoms that overlap with those of genitopatellar syndrome (described above), although some of the specific developmental abnormalities differ between the two conditions. Mutations that cause the SBBYS variant of Ohdo syndrome have been identified throughout the *KAT6B* gene, although many of them occur in exon 18. Studies suggest that these mutations likely prevent the production of functional histone acetyltransferase from one copy of the *KAT6B* gene in each

cell. A shortage of this enzyme impairs the regulation of various genes during early development. Because these mutations lead to a reduction in the enzyme, they are described as "loss-of-function." However, it is unclear how these changes lead to the specific features of the condition.

cancers

Genetic changes involving the *KAT6B* gene have been associated with certain types of cancer. These mutations are somatic, which means they are acquired during a person's lifetime and are present only in certain cells. The genetic changes are chromosomal rearrangements (translocations) that disrupt the region of chromosome 10 containing the *KAT6B* gene. Researchers have found a translocation that attaches this region of chromosome 10 to part of chromosome 16 in some people with a cancer of blood-forming cells called acute myeloid leukemia (AML). This translocation has also been identified in some people with therapy-related myelodysplastic syndrome, a blood disorder that can occur after a person has undergone chemotherapy for another form of cancer.

It is unclear how translocations involving the *KAT6B* gene are related to the development of cancer. These changes likely alter histone modification, which could prevent normal regulation of gene activity. Impaired gene regulation may contribute to the growth of cancers by allowing abnormal cells to grow and divide uncontrollably.

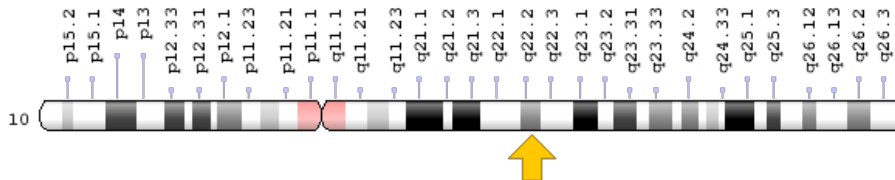
tumors

Somatic changes involving the *KAT6B* gene have also been identified in some people with uterine leiomyomas, which are noncancerous growths in the uterus that are also known as uterine fibroids. Uterine leiomyomas are common in adult women. These growths can cause pelvic pain and abnormal bleeding, and, in some cases, lead to an inability to have biological children (infertility). The genetic change associated with uterine leiomyomas is a translocation between the region of chromosome 10 containing the *KAT6B* gene and a particular region of chromosome 17. It is unclear how this translocation is related to tumor development. Changes in histone modification that impair normal gene regulation may allow certain cells to divide in an uncontrolled way, leading to the growth of a tumor.

Chromosomal Location

Cytogenetic Location: 10q22.2, which is the long (q) arm of chromosome 10 at position 22.2

Molecular Location: base pairs 74,824,927 to 75,032,623 on chromosome 10 (Homo sapiens Annotation Release 108, GRCh38.p7) (NCBI)



Credit: Genome Decoration Page/NCBI

Other Names for This Gene

- GTPTS
- histone acetyltransferase KAT6B
- histone acetyltransferase MORF
- histone acetyltransferase MOZ2
- histone acetyltransferase MYST4
- K(lysine) acetyltransferase 6B
- KAT6B_HUMAN
- monocytic leukemia zinc finger protein-related factor
- MORF
- MOZ-related factor
- MOZ2
- MYST-4
- MYST histone acetyltransferase (monocytic leukemia) 4
- MYST4
- qkf
- querkopf
- ZC2HC6B

Additional Information & Resources

Educational Resources

- Epigenomics Help (2010): Histone Modification "Writers"
https://www.ncbi.nlm.nih.gov/books/NBK45788/#epi_sci_bkgd.Histone_modification_write

GeneReviews

- KAT6B-Related Disorders
<https://www.ncbi.nlm.nih.gov/books/NBK114806>

Scientific Articles on PubMed

- PubMed
<https://www.ncbi.nlm.nih.gov/pubmed?term=%28KAT6B%5BTIAB%5D%29+OR+%28%28MORF%5BTIAB%5D%29+OR+%28MOZ2%5BTIAB%5D%29+OR+%28MYST4%5BTIAB%5D%29+OR+%28querkopf%5BTIAB%5D%29%29+AND+%28%28Genes%5BMH%5D%29+OR+%28Genetic+Phenomena%5BMH%5D%29%29+AND+english%5BIa%5D+AND+human%5Bmh%5D>

OMIM

- LEIOMYOMA, UTERINE
<http://omim.org/entry/150699>
- LEUKEMIA, ACUTE MYELOID
<http://omim.org/entry/601626>
- LYSINE ACETYLTRANSFERASE 6B
<http://omim.org/entry/605880>
- MYELODYSPLASTIC SYNDROME
<http://omim.org/entry/614286>

Research Resources

- Atlas of Genetics and Cytogenetics in Oncology and Haematology
<http://atlasgeneticsoncology.org/Genes/MYST4ID41488ch10q22.html>
- ClinVar
<https://www.ncbi.nlm.nih.gov/clinvar?term=KAT6B%5Bgene%5D>
- HGNC Gene Family: Lysine acetyltransferases
<http://www.genenames.org/cgi-bin/genefamilies/set/486>
- HGNC Gene Family: PHD finger proteins
<http://www.genenames.org/cgi-bin/genefamilies/set/88>
- HGNC Gene Family: Zinc fingers C2HC-type
<http://www.genenames.org/cgi-bin/genefamilies/set/66>

- HGNC Gene Symbol Report
http://www.genenames.org/cgi-bin/gene_symbol_report?q=data/hgnc_data.php&hgnc_id=17582
- NCBI Gene
<https://www.ncbi.nlm.nih.gov/gene/23522>
- UniProt
<http://www.uniprot.org/uniprot/Q8WYB5>

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Reviewed: February 2013
Published: March 21, 2017

Lister Hill National Center for Biomedical Communications
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National Institutes of Health
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